SEQUENCE LISTING

Rigel Pharmaceuticals, Incorporated

<120> TRAC1: Modulators of Lymphocyte Activation

:130: 021044-000600US

<140 > US 09/998,667 <141 > 2001-12-03

<150> US 60/282,432 <151> 2001-04-06

<160 > 18

<170> PatentIn Ver. 2.1

<210> 1 <211> 232 <212> PRT

<213> Homo sapiens

-: 220 ×

<223> human wild-type TRAC1 (FLJ20456)

<400> 1

Met Gly Ser Val Leu Ser Thr Asp Ser Gly Lys Ser Ala Pro Ala Ser 1 5 10 15

Ala Thr Ala Arg Ala Leu Glu Arg Arg Arg Asp Pro Glu Leu Pro Val

Thr Ser Phe Asp Cys Ala Val Cys Leu Glu Val Leu His Gln Pro Val 35 40 45

Arg Thr Arg Cys Gly His Val Phe Cys Arg Ser Cys Ile Ala Thr Ser 50 55 60

Leu Lys Asn Asn Lys Trp Thr Cys Pro Tyr Cys Arg Ala Tyr Leu Pro 65 70 75 80

Ser Glu Gly Val Pro Ala Thr Asp Val Ala Lys Arg Met Lys Ser Glu

Tyr Lys Asn Cys Ala Glu Cys Asp Thr Leu Val Cys Leu Ser Glu Met 100 105 110

Arg Ala His Ile Arg Thr Cys Gln Lys Tyr Ile Asp Lys Tyr Gly Pro 115 120 125

Leu Gln Glu Leu Glu Glu Thr Ala Ala Arg Cys Val Cys Pro Phe Cys 130 140

Glm Arg Glu Leu Tyr Glu Asp Ser Leu Leu Asp His Cys Ile Thr His 145 150 155 160

His Arg Ser Glu Arg Arg Pro Val Phe Cys Pro Leu Cys Arg Leu Ile 165 Pro Asp Glu Asn Pro Ser Ser Phe Ser Gly Asn Leu Ile Arg His Leu 185 Gln Val Ser His Thr Leu Phe Tyr Asp Asp Phe Ile Asp Phe Asn Ile 195 200 Ile Glu Glu Ala Leu Ile Arq Arq Val Leu Asp Arg Ser Leu Leu Glu 215 Tvr Val Asn His Ser Asn Thr Thr 230 <210> 2 <211> 2031 <212> DNA <213> Homo sapiens <220> <223> human wild-type TRAC1 cDNA (FLJ20456) <220> <221> CDS <222> (498)..(1196) <223> TRAC1 <400> 2 agcqqaqqtc atttttgcag cttattgtga tgacaacagt ggaggatggt cttccacttc 60 accttaaaag cggctgttct ctgattatca ttaagcatgg ccacgcccgc acttaacttc 120 tgacagtggg gaaagcagct gtgtgtgata gcttggaagg tttactgctg cctcaagtcc 180 tettetetge agttgaggtt teaggtttea atecteecaa taccacaaga cagageacgg 240 ggeggetgee geeteegeet eegegeetta acetaggegg ettgeegaag ateteageee 300 cgcggccgcg cgctcgccct gccctagacc agggttgggc gcagcggcgg aggtggcttc 360 tgggctgcgc gagctgggag agctgggagg cggcgatcgc agctgggccg ggacttcctt 420 cctccaccgc acggcaacaa aacaaccctg cggcaggcac tgagtgcttc gcagctgtct 480 gggcgagagg cacagegatg ggctccgtgc tgagcacega cageggcaaa teggegeeeg 540 octotgecae egegegggee etggagegea ggagggaeee ggagttgeee gteaegteet 600 togactgogc ogtgtgoctt gaggtgttac accagootgt coggaccogc tgcggccacg 660 tattotgoog ttootgtatt gotacoagto tgaagaacaa caagtggaco tgtoottatt 720 qeeqqqeata totteettea gaaggagtte cageaactga tgtagecaaa agaatgaaat 780 cagagtataa gaactgcgct gagtgtgaca ccctggtttg cctcagtgaa atgagggcac 840 atatteggae ttgteagaag tacatagata agtatggaee actacaagaa ettgaggaga 900 cagcagcaag gtgtgtatgt cocttttgtc agagggaact gtatgaagac agcttgctgg 960 atcattgtat tactcatcac agatcggaac ggaggcctgt gttctgtcca ctttgccgtt 1020 taatacccga tgagaatcca agcagcttca gcggcaattt aataagacat ctgcaagtta 1080 gtcacacttt gttttatgat gatttcatag attttaatat aattgaggaa gctcttatcc 1140 gaagagtett agaceggtea ettettgaat atgtgaatea etegaacace acataatttt 1200 attaaaacga agggaaaagg gaccactgaa ttgcaccatt taagatgctg cttgaacaaa 1260 tgggagggaa gttgtcaatg attgatgggc aaaaatgtac aacacagtta tgtgtttgtc 1320 catgittatt gitatagige attiaaaaac tgetttaatt tiaatggitt aaateigitt 1380 tacateetty agattettae asatetaaca asaaaaaaaa ttatetacat cagteattyt 1440 tacatggaaa agacaggtgg taggcaagta ggtggaggat ctcggtttgc aaattagata 1500 atactotgtg tataatgota catatoaata actacoatoa tggttaggoa cgataactaa 1560 totttgttot gtgtaaaaaa atatggagag tgaaacaaag tgcagacatt caaagaaata 1620 agaaatotgo tocaatgoto tigitotaat ototaatagg tiaaogitaa taatotigia 1680 tggqagttgg aaaggaaaat tttggaagtc aagaaagtcc atttaggccg gacgcggtgg 1740

ottacgettg tagteecage actttgggag gotgaggeag geggateaca gggtegggag 1800 ttegagaeca geetggeeaa eactggtete tgtgaaacte egtetetaet gaaaatgeaa 1860

agattggctg gacgtgttgg cgggcatctg tgataccagc tacttgggag gctgaggcag 1920 aagaatcgct tgagcccggg aggcggaggt tgcagtgagc tgagatcgcg ccagtacact 1980 ccagcotggg taaragagot agactocato traaaaaaaaa aaaaaaaaaa a <210> 3 <211> 1620 <212> DNA <213> Homo sapiens <220> <223> human wild-type TRAC1 shorter cDNA <220> <221> CDS <222> (383)..(1081) <223> TRAC1 <400>3acttetgaca gtggggaaag cagetgtgtg tgatagettg gaaggtttae tgetgeetca 60 agtectette tetgeagttg aggttteagg ttteaateet eecaatacea caagacagag 120 cacggggcgg etgecgeete egeeteegeg cettaaceta ggeggettge egaagatete 180 agccccgcgg ccgcgctc gccctgccct agaccagggt tgggcgcagc ggcggaggtg 240 gettetggge tgegegaget gggagagetg ggaggeggeg ategeagetg ggeegggaet 300 teetteetee acegeaegge aacaaaacaa eeetgeggea ggeaetgagt gettegeage 360 tgtctgggcg agaggcacag cgatgggctc cgtgctgagc accgacagcg gcaaatcggc 420 gcccgcctct gccaccgcgc gggccctgga gcgcaggagg gacccggagt tgcccgtcac 480 gteettegae tgegeegtgt geettgaggt gttacaccag cetgteegga eeegetgegg 540 ccacgtattc tgccgttcct gtattgctac cagtctgaag aacaacaagt ggacctgtcc 600 thattgoogg goatatotto ottoagaagg agttocagca actgatgtag ccaaaagaat 660 gaaatcagag tataagaact gegetgagtg tgacaccetg gtttgeetca gtgaaatgag 720 ggcacatatt cggacttgtc agaagtacat agataagtat ggaccactac aagaacttga 780 ggagacagca gcaaggtgtg tatgtccctt ttgtcagagg gaactgtatg aagacagctt 840 gctggatcat tgtattactc atcacagatc ggaacggagg cctgtgttct gtccactttg 900 ccgtttaata cccgatgaga atccaagcag cttcagcggc aatttaataa gacatctgca 960 agttagtcac actttgtttt atgatgattt catagatttt aatataattg aggaagctct 1020 tatccgaaga gtcttagacc ggtcacttct tgaatatgtg aatcactcga acaccacata 1080 attttattaa aacgaaggga aaagggacca ctgaattgca ccatttaaga tgctgcttga 1140 acaaatggga gggaagttgt caatgattga tgggcaaaaa tgtacaacac agttatgtgt 1200 ttgtccatgt ttattgttat agtgcattta aaaactgctt taattttaat ggtttaaatc 1260 tgttttacat ccttgagatt cttacacatc taacaacaaa aaaaattatc tacatcagtc 1320 attgttacat ggaaaagaca ggtggtaggc aagtaggtgg aggatctcgg tttgcaaatt 1380 agataatact ctgtgtataa tgctacatat caataactac catcatggtt aggcacgata 1440 actaatettt gttetgtgta aaaaaatatg gagagtgaaa caaagtgcag acattcaaag 1500 aaataagaaa totgotocaa tgotottgtt otaatotota ataggttaac gttaataato 1560 ttgtatggga gttggaaagg aaaattttgg aagtcaagaa agtccattta ggccggacgc 1620 <210> 4 <211> 765 <212> DNA :213> Hcmo sapiens :220> :223> cENA encoding truncated version of human TRAC1 ·:400> 4 geggetgeeg ceteegeete egegeettaa estaggegge ttgeegaaga teteageeee 60 geggeegege getegeeetg eectagaeea gegttgggeg eageggegga ggtggettet 120 gggctgcgcg agctgggaga gctgggaggc ggcgatcgca gctgggccgg gacttccttc 180

ctccaccgca cggcaacaaa acaaccetge ggcaggcact gagtgetteg cagetgtetg 240

```
ggcgagaggc acagcgatgg gctccgtgct gagcaccgac agcggcaaaat cggcgcccgc 300
ctctgccacc gcgcgggccc tggagcgcag gagggacccg gagttgcccg tcacgtcctt 360
cgactgcgcc gtgtgccttg aggtgttaca ccagcctgtc cggacccgct gcggccacgt 420
attoctgccgt tootgtattg ctaccagtot gaagaacaac aagtggacot gtoottattg 480
ccgggcatat cttccttcag aaggagttcc agcaactgat gtagccaaaa gaatgaaatc 540
agagtataag aactgegetg agtgtgacae eetggtttge etcagtgaaa tgagggeaca 600
tattcggact tgtcagaagt acatagataa gtatggacca ctacaagaac ttgaggagac 660
agcagcaagg tgtgtatgtc ccttttgtca gagggaactg tatgaagaca gcttgctgga 720
tcattgtatt actcatcaca gatcggaacg gaggcctgtg ttctg
<210> 5
<211> 4983
<212> DNA
<213> Homo sapiens
<220>
<223> human wild-type TRAC1 genomic sequence
<220>
<221> CDS
<222> (3634)..(4332)
<223> TRAC1
<400> 5
acttttagtt gaatttetga tteetgeeat caaaagtaag tettgeagga atttttagat 60
aatattaagg atttggaatt taatcctaag gattaggaga gctattaaag gattttgtgc 120
atggggtgac acaagatgtt tgcttttcaa aagatcactt tagttgccat gtggataata 180
aactggagag aggcaatgat ggatgcgggt agagcagtta ggaactactg ccattaagtc 240
acacaagaga atgcagtgat ttaaaataag cggtggctat ggaaatagaa gaaaatgagc 300
atggtggctc acgcctgtaa tcccagcact ttgggaggcc aaggcaggcg gatcatgagg 360
teaggagtte acgaetagee tggccaacat ggtgaaacte egtetetace aaaaatataa 420
taaaattagc cgggtgtggt ggcatgcgcc tgcaatccca gctactcagg aggctgaggc 480
aggagaatca cttgaaccca ggaggcagag gttgcagtga gctgagattg cgccactgca 540
cacttccage etgggegaca gagagagaet teateteaga aaaaaaagaa etaetgagat 600
acatattgga ggcaggattg tgattatgct tgattcaatg tgaatgagga agaggaagaa 660
atcaaggatg acttccaggt gtctagactg agctataaag tggatcatag tgccatttgc 720
taaaagagag atcaaccact ggaggaggct gctactataa tgagttcatt attggagaca 780
ttgggctgag ggtgtttatg tcaaatggtc agtcatgtaa gctattaggc tattggacat 840
ttgaatggcc tggggtagag ataaagatgt gaaagttttt ggaaccgaaa tagtgacaga 900
ggtcctaaga cagaatccta ggatctccag cccagagcca gtggagaagg agactctggc 1020
aagggagaag aaacaatagc cagaaagaaa gagggaaagt cagagtgttt ccagaaggag 1080
gggatggtca qcactaacaa acatagttga ggggtcaagc aaaaaaaaata gctgaaaaga 1140
atctattqqa attaqttaca tgaacgtcac cagtgacact gataataaag cagtttttgg 1200
acagatggag gtggagaggt tggttcagaa cccagactgg actgaataag aagtgaataa 1260
ttaagaaatg atgacaaaat gtagaggatc agatcaagag atttggcctt gaagcggata 1320
tggggcagga gttgaggcat aagtgggatg aaagggaggt ttttgttttc cttttgaaga 1380
tgggaataac taccttttca tottatttcc coccoccac cgcctgcccc ccaccacca 1440
tgctccaget ccataggtet treetttett tggttetgee teatreactg treecagtee 1500
acgettttte etaaatettt gaatggetea ettetttea ttatteatgg etetgeteaa 1560
atgleactic atccaaaaat tettetetga gettattett etecteatig ceatttactg 1620
ttttatette tteatggete tgattatetg aaatgatttt gtteatttgt ttatggattg 1680
attacatgta tetteetttt gittigtigtt igittittigag acagagiete geteigtiae 1740
ccacgatgga gtgtagtggc atggtctcgg stcactgcaa cetesacete ttgggtgcaa 1800
gtggttctcc tgcctcagcc tcccaagtag stgggactac acgcatgtgc caccacacct 1860
ggctaatttt tgtattttta gtacaaaaat ttttactaaa gaaatgtatt tctagtagag 1920
acggggtttc actatgttgg caaggctggt attgaactcc tgacattgtg atccacctgc 1980
ctoggoctoc caaattgotg ggattteagg ogtgageeac tgeacecage etgttttggt 2040
ttttttgagad agggtetede tetgttgede aggetggagt geagtggeat gatetegget 2100
cactgoagec tecacettic eggiteaage gattetigtg ceteagecte ceaagtaget 2160
```

cacacccage taagttttgt attittagta gagacagggt ticaccatgt tggccagget 2220 ggtctcaaac teetgaeete aggtgatetg teettettge eteccaaagt getgggatta 2280 caggogtgag ccactgoocc ggottggttt gttttttgag acatggtotc actotgtogc 2340 acaggacgga gtgcagtagt gtgatcttgg ctcccgtgct caagtgatcc tcccacctca 2400 gcctcccaaa gtgctgaaat tacaggcgtg agccacggta cccggcctat atctcctaat 2460 taaaagtaaa gootgggtga agtttatggo otgotaaata ogtacatgta ttatatgogt 2520 tattatatcc catgtgttat attatatatt attgtatatg tgtattatat actatgtact 2580 attottacaa otgtatgtga atotataatt atotoaaagt tittgttittg tittttaaag 2640 taagttetgt gagaactggg attttgtttg ettgttgaeg attgtgtett eateactaga 2700 accgagtctg gcgtgcagtg ggcactaaat aagtcttcgt aaaaagtgta aattaacgcc 2760 eggetaattt ttgtattatt ageagegaeg gggttteace atgttggeea ggetggtete 2820 gaacteetge ceteaageag gagttteaag eeegeetegg eeteeeaaag tgttggaatt 2880 acaggegtgg gecacegege etggetgate atgtttaaag ggeggggtgg ggagegtaca 2940 aacaacagag gaaatcctga gccgcagagg aaactggaga tggcagggtt tagcactaga 3000 atctctgtag gagaggtaag attggaatcc taatctccag gactctttcc actacccagg 3060 ctatetetee attaatggae tattgattgg atgacagagt agegatagea eegtettagg 3120 agacgcccaa tcataggtca taggtcattt ttgcagctta ttgtgatgac aacagtggag 3180 gatggtcttc cacttcacct taaaagcggc tgttctctga ttatcattaa gcatggccac 3240 qcccqcactt aacttctgac agtggggaaa gcagctgtgt gtgatagctt ggaaggttta 3300 ctgctgcctc aagtcctctt ctctgcagtt gaggtttcag gtttcaatcc tcccaatacc 3360 acaagacaga gcacggggcg gctgccgcct ccgcctccgc gccttaacct aggcggcttg 3420 ccgaagatet cageceegeg geegegeget egeeetgeee tagaceaggg ttgggegeag 3480 cggcggaggt ggcttctggg ctgcgcgagc tgggagagct gggaggcggc gatcgcagct 3540 gggccgggac ttccttcctc caccgcacgg caacaaaaca accctgcggc aggcactgag 3600 tgcttcgcag ctgtctgggc gagaggcaca gcgatgggct ccgtgctgag caccgacagc 3660 ggcaaatcgg cgcccgcctc tgccaccgcg cgggccctgg agcgcaggag ggacccggag 3720 ttgcccgtca cgtccttcga ctgcgccgtg tgccttgagg tgttacacca gcctgtccgg 3780 accogotgog gocacgtatt otgoogttoo tgtattgota coagtotaaa gaacaacaag 3840 tggacctgtc cttattgccg ggcatatctt ccttcagaag gagttccagc aactgatgta 3900 gccaaaagaa tgaaatcaga gtataagaac tgcgctgagt gtgacaccct ggtttgcctc 3960 agtgaaatga gggcacatat tcggacttgt cagaagtaca tagataagta tggaccacta 4020 caagaacttg aggagacagc agcaaggtgt gtatgtccct tttgtcagag ggaactgtat 4080 gaagacagct tgctggatca ttgtattact catcacagat cggaacggag gcctgtgttc 4140 tgtccacttt gccgtttaat acccgatgag aatccaagca gcttcagtgg cagtttaata 4200 agacatetge aagttagtea caetttgttt tatgatgatt teatagattt taatataatt 4260 gaggaagete ttateegaag agtettagae eggteaette ttgaatatgt gaateaeteg 4320 aacaccacat aattttatta aaacgaaggg aaaagggacc actgaattgc accatttaag 4380 atgctgcttg aacaaatggg agggaagttg tcaatgattg atgggcaaaa atgtacaaca 4440 cagttatgtg tttgtccatg tttattgtta tagtgcattt aaaaactgct ttaattttaa 4500 tggtttaaat ctgttttaca tccttgagat tcttacacat ctaacaacaa aaaaaattat 4560 ctacatcagt cattgttaca tggaaaagac aggtggtagg caagtaggtg gaggatctcg 4620 gtttgcaaat tagataatac tctgtgtata atgctacata tcaataacta ccatcatggt 4680 taggcacgat aactaatett tgttetgtgt aaaaaaatat ggagagtgaa acaaagtgca 4740 gacattcaaa gaaataagaa atctgctcca atgctcttgt tctaatctct aataggttaa 4800 cgttaataat cttgtatggg agttggaaag gaaaattttg gaagtcaaga aagtccattt 4860 aggccggacg cggtggctta cgcttgtaat cccagcactt tgggaggctg aagcaggcgg 4920 atcacaaggt caggagttcg agaccagcct ggccaacact ggtctctgtg aaactccgtc 4980 tct

```
<210> 6
<211> 1408
<212> DNA
<213> Mus sp.
<220>
<223> mouse TRAC1 cDNA
```

```
<220>
<221> modified base
<222> (167)..(168)
<213> n = a, g, c or t
<220>
<221> CDS
<222> (159)..(869)
<223> TRAC1
<400> 6
tgegegeget eegeetgege tecacegaga ggeeeegegg egeggetgge egageeagga 60
ggcggcgatc cggcctgggc cgggacttcc ttcctcccgg gcgggacaac agaaccaccc 120
gcagccagca gggagctcct ggcagctcgc tgagctagag cgcaagnntg ggctccctgc 180
tgagcagcga cagetecaag teegegeeeg eeteegeeae eeegeggaet etggagegea 240
geggggaete ggagetgeee ateaceteet tegactgete agtgtgtetg gaggtgetae 300
accagooggt coggacoogc tgtggccacg tgttctgccg atcttgcatt gcgaccagta 360
taaagaacaa taataaatgg acatgtccat actgccgggc ataccttcct tcagagggag 420
tgcccgcaac tgacatagcc aagaggatga agtcagaata ccagaactgt gctgagtgtg 480
gaactotggt ttgcctcagt gacatgaggg cgcacataag gacctgtgag aagtacatcg 540
ataaatatgg cccgctgcta gaacttggcg acaccacagc aagatgtgta tgtccatttt 600
gtcagcggga actggatgaa gactgcttgc tggatcattg cattatccac cacagatcag 660
anaggaggcc cgtgttctgt ccactttgcc attcacgacc tgatgaaagc ccaagtacct 720
tcaatggcag tttaattaga catttgcaag tcagtcacac tttgttttat gatgatttca 780
tagattttga tataattgag gaagccatta ttcgcagagt gctagaccgc tcacttcttg 840
aatatgtgaa tcagtcaaac accacataat tttatgacta ggaaggggac cattcactcg 900
taccatttaa gatgctgctt gaaggactgg agggggattg tcgacgtttg atggagaaac 960
atgtactaca gtattacctg cttgtctttg ttatgttgca ttcaaaaacc gtgttcattc 1020
tggcttagat ctgcccttac attcttgagt ctattagaca tttaaccacg agacacagcc 1080
tctcagccat taacagatgg aagagacacg acacaggctg gctatgtgat ggagcagctt 1140
cctgtgtcca ttagtgatgt gtgtataacg ctacatattc ccaactgtca tcatggttag 1200
qcqaqaaaqc caatctctgt tccatgctaa aagacggaga aaggaacaaa atatggacat 1260
taarqqaaat ctgaaatgta ttcaaattct catgctctaa tctccaataa ggaacgtgaa 1320
taatottatg aaaaggagga aagggaagaa titgaagtoa ccaaaatoca atttagocaa 1380
attatagtac ataatataat actgcagc
<210> 7
<211> 239
<212> PRT
<213> Mus sp.
%223> mouse TRAC1 protein (3rd frame)
<220>
<221> MOD RES
<222> (3)
<223> Xaa = Arg or Ser
< 220>
<221> MOD RES
<222> (4)
<223> Xaa = Met, Val or Leu
<400> 7
Ser Ala Xaa Xaa Gly Ser Leu Leu Ser Ser Asp Ser Ser Lys Ser Ala
                                     10
Pro Ala Ser Ala Thr Pro Arg Thr Leu Glu Arg Ser Gly Asp Ser Glu
             20
                                 25
```

Leu Pro Ile Thr Ser Phe Asp Cys Ser Val Cys Leu Glu Val Leu His

Gln Pro Val Arg Thr Arg Cys Gly His Val Phe Cys Arg Ser Cys Ile 50  $\,$  50

Ala Thr Ser Ile Lys Asn Asn Asn Lys Trp Thr Cys Pro Tyr Cys Arg 65 70 75 80

Ala Tyr Leu Pro Ser Glu Gly Val Pro Ala Thr Asp Ile Ala Lys Arg 85 90 95

Met Lys Ser Glu Tyr Gln Asn Cys Ala Glu Cys Gly Thr Leu Val Cys 100 105 110

Leu Ser Asp Met Arg Ala His Ile Arg Thr Cys Glu Lys Tyr Ile Asp 115 120 125

Lys Tyr Gly Pro Leu Leu Glu Leu Gly Asp Thr Thr Ala Arg Cys Val 130 135 140

Cys Pro Phe Cys Gln Arg Glu Leu Asp Glu Asp Cys Leu Leu Asp His 145 150 155 160

Cys Ile Ile His His Arg Ser Glu Arg Arg Pro Val Phe Cys Pro Leu 165 170 175

Cys His Ser Arg Pro Asp Glu Ser Pro Ser Thr Phe Asn Gly Ser Leu 180 185 190

Ile Arg His Leu Gln Val Ser His Thr Leu Phe Tyr Asp Asp Phe Ile 195 200 205

Asp Phe Asp Ile Ile Glu Glu Ala Ile Ile Arg Arg Val Leu Asp Arg 210 215 220

Ser Leu Leu Glu Tyr Val Asn Gln Ser Asn Thr Thr Phe Tyr Asp 225 230 235

<210> 8

<111> 228

<212> PRT

<213> Homo sapiens

<220>

<223> znf313 sequence with ring domain

<400> 8

Met Ala Ala Gln Gln Arg Asp Cys Gly Gly Ala Ala Gln Leu Ala Gly
1 5 10 15

Pro Ala Ala Glu Ala Asp Pro Leu Gly Arg Phe Thr Cys Pro Val Cys

Leu Glu Val Tyr Glu Lys Pro Val Gln Val Pro Cys Gly His Val Phe 35 40 45

Cys Ser Ala Cys Leu Gln Glu Cys Leu Lys Pro Lys Lys Pro Val Cys 50 55 60

Gly Val Cys Arg Ser Ala Leu Ala Pro Gly Val Arg Ala Val Glu Leu 65 70 75 80

Glu Arg Gln Ile Glu Ser Thr Glu Thr Ser Cys His Gly Cys Arg Lys 85 90 95

Asn Phe Phe Leu Ser Lys Ile Arg Ser His Val Ala Thr Cys Ser Lys 100 105 110

Tyr Gln Asn Tyr Ile Met Glu Gly Val Lys Ala Thr Ile Lys Asp Ala 115 120 125

Ser Leu Gln Pro Arg Asn Val Pro Asn Arg Tyr Thr Phe Pro Cys Pro 130 135 140

Tyr Cys Pro Glu Lys Asn Phe Asp Gln Glu Gly Leu Val Glu His Cys 145 150 155 160

Lws Leu Phe His Ser Thr Asp Thr Lys Ser Val Val Cys Pro Ile Cys 165 170 175

Ala Ser Met Pro Trp Gly Asp Pro Asn Tyr Arg Ser Ala Asn Phe Arg 180 185 190

Glu His Ile Gln Arg Arg His Arg Phe Ser Tyr Asp Thr Phe Val Asp 195 200 205

Tyr Asp Val Asp Glu Glu Asp Met Met Asn Gln Val Leu Gln Arg Ser 210 215 220

Ile Ile Asp Gln 205

<210> 9

<211> 245

<212> PRT

<213> Homo sapiens

<220>

<223> STRIN sequence with righ domain

<400 > 9

Met Ala Glu Asp Leu Ser Ala Ala Thr Ser Tyr Thr Glu Asp Asp Phe

1 5 10 15

Tyr Cys Pro Val Cys Gln Glu Val Leu Lys Thr Pro Val Arg Thr Thr 20 25 30

Ala Cys Gln His Val Phe Cys Arg Lys Cys Phe Leu Thr Ala Met Arg 35 40 45

Glu Ser Gly Ala His Cys Pro Leu Cys Arg Gly Asn Val Thr Arg Arg 50 55 60

Glu Arg Ala Cys Pro Glu Arg Ala Leu Asp Leu Glu Asn Ile Met Arg 65 70 75 80

Lys Phe Ser Gly Ser Cys Arg Cys Cys Ala Lys Gln Ile Lys Phe Tyr 85 90 95 Arg Met Arg His His Tyr Lys Ser Cys Lys Lys Tyr Gln Asp Glu Tyr
100 105 110

Gly Val Ser Ser Ile Val Pro Asn Phe Gln Ile Ser Gln Asp Ser Val 115 120 125

Gly Asn Ser Asn Arg Ser Glu Thr Ser Thr Ser Asp Asn Thr Glu Thr
130 135 140

Tyr Gln Glu Asn Thr Ser Ser Ser Gly His Pro Thr Phe Lys Cys Pro 145 150 155 160

Leu Cys Gln Glu Ser Asn Phe Thr Arg Gln Arg Leu Leu Asp His Cys 165 170 175

Asn Ser Asn His Leu Phe Gln Ile Val Pro Val Thr Cys Pro Ile Cys 180 185 190

Val Ser Leu Pro Trp Gly Asp Pro Ser Gln Ile Thr Arg Asn Phe Val 195 200 205

Ser His Leu Asn Gln Arg Arg Gln Phe Asp Tyr Gly Glu Phe Val Asn 210 215 220

Leu Gln Leu Asp Glu Glu Thr Gln Tyr Gln Thr Ala Val Glu Glu Ser 225 230 235 235

Phe Gln Val Asn Ile

<210> 10

<211> 50

<212> PRT

<213> Artificial Sequence

<220>

<400> 10

Val Thr Ser Phe Asp Cys Ala Val Cys Leu Glu Val Leu His Gln Pro

Val Arg Thr Arg Cys Gly His Val Phe Cys Arg Ser Cys Ile Ala Thr 20 25 30

Ser Leu Lys Asn Asn Lys Trp Thr Cys Pro Tyr Cys Arg Ala Tyr Leu 35 40 45

Pro Ser 50

<210> 11

:2112 50

<212> PRT

<213> Artificial Sequence

<220> <223> Description of Artificial Sequence:human znf313 ring finger domain <400> 11 Leu Gly Arg Phe Thr Cys Pro Val Cys Leu Glu Val Tyr Glu Lys Pro 1.0 Tal Gln Val Pro Cys Gly His Val Phe Cys Ser Ala Cys Leu Gln Glu 20 Cys Leu Lys Pro Lys Lys Pro Val Cys Gly Val Cys Arg Ser Ala Leu 40 Ala Pro 50 <210> 12 <211> 50 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence:human STRIN ring finger domain <400> 12 Glu Asp Asp Phe Tyr Cys Pro Val Cys Gln Glu Val Leu Lys Thr Pro 5 10 Yal Arg Thr Thr Ala Cys Gln His Val Phe Cys Arg Lys Cys Phe Leu 20 25 Thr Ala Met Arg Glu Ser Gly Ala His Cys Pro Leu Cys Arg Gly Asn Val Thr 50 <210> 13 <211> 50 <212> PRT 3213: Artificial Sequence <223> Description of Artificial Sequence:human TRAF6 ring finger domain -:400> 13 Glu Ser Lys Tyr Glu Cys Pro Ile Cys Leu Met Ala Leu Arg Glu Ala 5 Val Gln Thr Pro Cys Gly His Arg Phe Cys Lys Ala Cys Ile Ile Lys

Ser Ile Arg Asp Ala Gly His Lys Cys Pro Val Asp Asn Glu Ile Leu 35 40 45

```
Leu Glu
    50
<210> 14
<211> 50
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:human c-Cbl
      ring finger domain
<400: 14
Ser Thr Phe Gln Leu Cys Lys Ile Cys Ala Glu Asn Asp Lys Asp Val
                       10
Lys !le Glu Pro Cys Gly His Leu Met Cys Thr Ser Cys Leu Thr Ser
Trp Gln Glu Ser Glu Gly Gln Gly Cys Pro Phe Cys Arg Cys Glu Ile
                            40
Lys Gly
    50
<210> 15
<211> 50
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence:human BRCA1
     ring finger domain
<400: 15
Leu Glu Cys Pro Ile Cys Leu Glu Leu Ile Lys Glu Pro Val Ser Thr
Lys Cys Asp His Ile Phe Cys Lys Phe Cys Met Leu Lys Leu Leu Asn
                                 25
Gln Lys Lys Gly Pro Ser Gln Cys Pro Leu Cys Lys Asn Asp Ile Thr
                            40
Lys Arg
     50
<210:- 16
<211> 50
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence:human BAR ring
      finger domain
```

.

-:400> 16 Val Ser Glu Phe Ser Cys His Cys Cys Tyr Asp Ile Leu Val Asn Pro 1.0 Thr Thr Leu Asn Cys Gly His Ser Phe Cys Arg His Cys Leu Ala Leu 25 30 20 Trp Trp Ala Ser Ser Lys Lys Thr Glu Cys Pro Glu Cys Arg Glu Lys 40 Trp Glu 50 <210: 17 <211> 49 -::11: PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence:human RAG1 ring finger domain <400> 17 Lys Ser Ile Ser Cys Gln Ile Cys Glu His Ile Leu Ala Asp Pro Val Glu Thr Asn Cys Lys His Val Phe Cys Arg Val Cys Ile Leu Arg Cys Leu Lys Val Met Gly Ser Tyr Cys Pro Ser Cys Arg Tyr Pro Cys Phe 40 Pro H210H 18 <211> 200 <212> PRT 1213: Artificial Sequence <223> Description of Artificial Sequence: flexible linker 220 -<221> MOD\_RES <2225 (6)..(200) <223> Gly at positions 6-200 may be present or absent <400> 18 

3.5

 31y
 31y</td